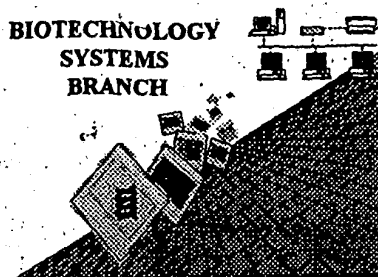


## RAW SEQUENCE LISTING ERROR REPORT

BIOTECHNOLOGY  
SYSTEMS  
BRANCH



0400  
04/13/2001

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/808,832

Source: OIPE

Date Processed by STIC: 3/30/2001

**THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.**

**PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:**

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,**
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY**

**FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.**

**FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.**

**PATENTIN 2.1 e-mail help: [patin21help@uspto.gov](mailto:patin21help@uspto.gov) or phone 703-306-4119 (R. Wax)**

**PATENTIN 3.0 e-mail help: [patin3help@uspto.gov](mailto:patin3help@uspto.gov) or phone 703-306-4119 (R. Wax)**

**TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:**

### **Checker Version 3.0**

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 - 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

<http://www.uspto.gov/web/offices/pac/checker>

# Raw Sequence Listing Error Summary

## ERROR DETECTED SUGGESTED CORRECTION

SERIAL NUMBER: 09/808,832

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1        Wrapped Nucleics      The number/text at the end of each line "wrapped" down to the next line.  
This may occur if your file was retrieved in a word processor after creating it.  
Please adjust your right margin to .3, as this will prevent "wrapping".
- 2        Wrapped Aminos      The amino acid number/text at the end of each line "wrapped " down to the next line.  
This may occur if your file was retrieved in a word processor after creating it.  
Please adjust your right margin to .3, as this will prevent "wrapping".
- 3        Incorrect Line Length      The rules require that a line not exceed 72 characters in length. This includes spaces.
- 4        Misaligned Amino Acid      The numbering under each 5th amino acid is misaligned. This may be caused by the use of tabs  
Numbering      between the numbering. It is recommended to delete any tabs and use spacing between the numbers.
- 5        Non-ASCII      This file was not saved in ASCII (DOS) text, as required by the Sequence Rules.  
Please ensure your subsequent submission is saved in ASCII text so that it can be processed.
- 6        Variable Length      Sequence(s)        contain n's or Xaa's which represented more than one residue.  
As per the rules, each n or Xaa can only represent a single residue.  
Please present the maximum number of each residue having variable length and  
indicate in the (ix) feature section that some may be missing.
- 7        PatentIn ver. 2.0 "bug"      A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid  
sequence(s)       . Normally, PatentIn would automatically generate this section from the  
previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section  
to the subsequent amino acid sequence. **This applies primarily to the mandatory <220>-<223>**  
**sections for Artificial or Unknown sequences.**
- 8        Skipped Sequences      Sequence(s)        missing. If intentional, please use the following format for each skipped sequence:  
(OLD RULES)      **(2) INFORMATION FOR SEQ ID NO:X:**  
                                 (i) **SEQUENCE CHARACTERISTICS:**(Do not insert any headings under "SEQUENCE CHARACTERISTICS")  
                                 (xi) **SEQUENCE DESCRIPTION:SEQ ID NO:X:**  
                                 **This sequence is intentionally skipped**  
  
Please also adjust the "(iii) NUMBER OF SEQUENCES:" response to include the skipped sequence(s).
- 9        Skipped Sequences      Sequence(s)        missing. If intentional, please use the following format for each skipped sequence.  
(NEW RULES)      **<210> sequence id number**  
                                 **<400> sequence id number**  
                                 **000**
- 10        Use of n's or Xaa's      Use of n's and/or Xaa's have been detected in the Sequence Listing.  
(NEW RULES)      Use of <220> to <223> is MANDATORY if n's or Xaa's are present.  
                                 In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
- 11        Use of <213>Organism      Sequence(s)        are missing this mandatory field or its response.  
(NEW RULES)
- 12        Use of <220>Feature      Sequence(s)        are missing the <220>Feature and associated headings.  
(NEW RULES)      Use of <220> to <223> is MANDATORY if <213>ORGANISM is "Artificial Sequence" or "Unknown"  
                                 Please explain source of genetic material in <220> to <223> section.  
                                 (See "Federal Register," 6/01/98, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of new Rules)
- 13        PatentIn ver. 2.0 "bug"      Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted  
file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing).  
Instead, please use "File Manager" or any other means to copy file to floppy disk.

OIPE

RAW SEQUENCE LISTING  
 PATENT APPLICATION: US/09/808,832

DATE: 03/30/2001  
 TIME: 15:15:26

Input Set : A:\PTO.txt  
 Output Set: N:\CRF3\03302001\I808832.raw

Does Not Comply  
 Corrected Diskette Needed

3 <110> APPLICANT: DuPont Pharmaceuticals Company  
 5 <120> TITLE OF INVENTION: Peptidase-cleavable, targeted antineoplastic drugs and their therapeutic

6 use  
 8 <130> FILE REFERENCE: PH-7134  
 C--> 10 <140> CURRENT APPLICATION NUMBER: US/09/808,832  
 C--> 10 <141> CURRENT FILING DATE: 2001-03-15  
 10 <150> PRIOR APPLICATION NUMBER: 60/189,387  
 11 <151> PRIOR FILING DATE: 2000-03-15  
 13 <160> NUMBER OF SEQ ID NOS: 210  
 15 <170> SOFTWARE: PatentIn version 3.0  
 17 <210> SEQ ID NO: 1

18 <211> LENGTH: 5  
 19 <212> TYPE: PRT  
 20 <213> ORGANISM: Artificial  
 22 <220> FEATURE:  
 23 <223> OTHER INFORMATION: no comment  
 25 <220> FEATURE:  
 26 <221> NAME/KEY: MOD\_RES  
 27 <222> LOCATION: (1)..(1)  
 28 <223> OTHER INFORMATION: 4-methoxy-benzenesulfonyl-beta-alanine  
 31 <220> FEATURE:  
 32 <221> NAME/KEY: MOD\_RES  
 33 <222> LOCATION: (3)..(3)  
 34 <223> OTHER INFORMATION: homophenylalanine  
 37 <400> SEQUENCE: 1

39 Xaa Gly Xaa Tyr Leu  
 40 1 5

42 <210> SEQ ID NO: 2  
 43 <211> LENGTH: 5  
 44 <212> TYPE: PRT  
 45 <213> ORGANISM: Artificial  
 47 <220> FEATURE:  
 48 <223> OTHER INFORMATION: no comment  
 50 <220> FEATURE:  
 51 <221> NAME/KEY: MOD\_RES  
 52 <222> LOCATION: (1)..(1)  
 53 <223> OTHER INFORMATION: 1,2-C6H4(CO)2-histidine  
 56 <220> FEATURE:  
 57 <221> NAME/KEY: MOD\_RES  
 58 <222> LOCATION: (3)..(3)  
 59 <223> OTHER INFORMATION: homophenylalanine  
 62 <400> SEQUENCE: 2

64 Xaa Gly Xaa Tyr Leu  
 65 1 5  
 67 <210> SEQ ID NO: 3  
 68 <211> LENGTH: 5  
 69 <212> TYPE: PRT

*per 1.823 of Sequence Rules, valid response is Artificial Sequence*  
*give source of genetic material - see item 12 on Error Summary sheet*

## RAW SEQUENCE LISTING

DATE: 03/30/2001

PATENT APPLICATION: US/09/808,832

TIME: 15:15:26

Input Set : A:\PTO.txt

Output Set: N:\CRF3\03302001\I808832.raw

70 <213> ORGANISM: Artificial  
72 <220> FEATURE:  
73 <223> OTHER INFORMATION: no comment  
75 <220> FEATURE:  
76 <221> NAME/KEY: MOD\_RES  
77 <222> LOCATION: (1)..(1)  
78 <223> OTHER INFORMATION: acetyl-proline  
81 <400> SEQUENCE: 3  
OK 83 Xaa Leu Gly Leu Leu  
84 1 5  
86 <210> SEQ ID NO: 4  
87 <211> LENGTH: 5  
88 <212> TYPE: PRT  
89 <213> ORGANISM: Artificial  
91 <220> FEATURE:  
92 <223> OTHER INFORMATION: no comment  
94 <220> FEATURE:  
95 <221> NAME/KEY: MOD\_RES  
96 <222> LOCATION: (1)..(1)  
97 <223> OTHER INFORMATION: acetyl-proline  
100 <400> SEQUENCE: 4  
OK 102 Xaa Leu Gly Leu Leu  
103 1 5  
105 <210> SEQ ID NO: 5  
106 <211> LENGTH: 5  
107 <212> TYPE: PRT  
108 <213> ORGANISM: Artificial  
110 <220> FEATURE:  
111 <223> OTHER INFORMATION: no comment  
113 <220> FEATURE:  
114 <221> NAME/KEY: MOD\_RES  
115 <222> LOCATION: (2)..(2)  
116 <223> OTHER INFORMATION: beta alanine  
119 <220> FEATURE:  
120 <221> NAME/KEY: MOD\_RES  
121 <222> LOCATION: (1)..(1)  
122 <223> OTHER INFORMATION: acetyl-proline  
125 <400> SEQUENCE: 5  
OK 127 Xaa Xaa Gly Leu Leu  
128 1 5  
130 <210> SEQ ID NO: 6  
131 <211> LENGTH: 5  
132 <212> TYPE: PRT  
133 <213> ORGANISM: Artificial  
135 <220> FEATURE:  
136 <223> OTHER INFORMATION: no comment  
138 <220> FEATURE:  
139 <221> NAME/KEY: MOD\_RES  
140 <222> LOCATION: (2)..(2)

## RAW SEQUENCE LISTING

DATE: 03/30/2001

PATENT APPLICATION: US/09/808,832

TIME: 15:15:26

Input Set : A:\PTO.txt

Output Set: N:\CRF3\03302001\I808832.raw

141 &lt;223&gt; OTHER INFORMATION: 4-aminobutyric acid

144 &lt;220&gt; FEATURE:

145 &lt;221&gt; NAME/KEY: MOD\_RES

146 &lt;222&gt; LOCATION: (1)..(1)

147 &lt;223&gt; OTHER INFORMATION: acetyl-proline

150 &lt;400&gt; SEQUENCE: 6

OK 152 Xaa Xaa Gly Leu Leu

153 1 5

155 &lt;210&gt; SEQ ID NO: 7

156 &lt;211&gt; LENGTH: 5

157 &lt;212&gt; TYPE: PRT

158 &lt;213&gt; ORGANISM: Artificial

160 &lt;220&gt; FEATURE:

161 &lt;223&gt; OTHER INFORMATION: no comment

163 &lt;220&gt; FEATURE:

164 &lt;221&gt; NAME/KEY: MOD\_RES

165 &lt;222&gt; LOCATION: (2)..(2)

166 &lt;223&gt; OTHER INFORMATION: cyclohexylalanine

169 &lt;220&gt; FEATURE:

170 &lt;221&gt; NAME/KEY: MOD\_RES

171 &lt;222&gt; LOCATION: (1)..(1)

172 &lt;223&gt; OTHER INFORMATION: acetyl-proline

175 &lt;400&gt; SEQUENCE: 7

OK 177 Xaa Xaa Gly Leu Leu

178 1 5

180 &lt;210&gt; SEQ ID NO: 8

181 &lt;211&gt; LENGTH: 5

182 &lt;212&gt; TYPE: PRT

183 &lt;213&gt; ORGANISM: Artificial

185 &lt;220&gt; FEATURE:

186 &lt;223&gt; OTHER INFORMATION: no comment

188 &lt;400&gt; SEQUENCE: 8

190 Pro Leu Gly Leu Leu

191 1 5

193 &lt;210&gt; SEQ ID NO: 9

194 &lt;211&gt; LENGTH: 5

195 &lt;212&gt; TYPE: PRT

196 &lt;213&gt; ORGANISM: Artificial

198 &lt;220&gt; FEATURE:

199 &lt;223&gt; OTHER INFORMATION: no comment

201 &lt;220&gt; FEATURE:

202 &lt;221&gt; NAME/KEY: MOD\_RES

203 &lt;222&gt; LOCATION: (1)..(1)

204 <223> OTHER INFORMATION: MeOCH<sub>2</sub>CH<sub>2</sub>OCH<sub>2</sub>(=O)-proline

207 &lt;400&gt; SEQUENCE: 9

OK 209 Xaa Leu Gly Leu Leu

210 1 5

212 &lt;210&gt; SEQ ID NO: 10

213 &lt;211&gt; LENGTH: 5

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/09/808,832

DATE: 03/30/2001  
TIME: 15:15:26

Input Set : A:\PTO.txt  
Output Set: N:\CRF3\03302001\I808832.raw

214 <212> TYPE: PRT  
215 <213> ORGANISM: Artificial  
217 <220> FEATURE:  
218 <223> OTHER INFORMATION: no comment  
220 <220> FEATURE:  
221 <221> NAME/KEY: MOD\_RES  
222 <222> LOCATION: (1)..(1)  
223 <223> OTHER INFORMATION: MeOCH<sub>2</sub>CH<sub>2</sub>OCH<sub>2</sub>CH<sub>2</sub>OCH<sub>2</sub>C(=O)-proline  
226 <400> SEQUENCE: 10  
wt 228 Xaa Leu Gly Leu Leu  
229 1 5  
231 <210> SEQ ID NO: 11  
232 <211> LENGTH: 5  
233 <212> TYPE: PRT  
234 <213> ORGANISM: Artificial  
236 <220> FEATURE:  
237 <223> OTHER INFORMATION: no comment  
239 <220> FEATURE:  
240 <221> NAME/KEY: MOD\_RES  
241 <222> LOCATION: (1)..(1)  
242 <223> OTHER INFORMATION: H<sub>2</sub>NCH<sub>2</sub>CH<sub>2</sub>N(CH<sub>2</sub>CH<sub>2</sub>)<sub>2</sub>NCH<sub>2</sub>C(=O)-proline  
245 <400> SEQUENCE: 11  
wt 247 Xaa Leu Gly Leu Leu  
248 1 5  
250 <210> SEQ ID NO: 12  
251 <211> LENGTH: 5  
252 <212> TYPE: PRT  
253 <213> ORGANISM: Artificial  
255 <220> FEATURE:  
256 <223> OTHER INFORMATION: no comment  
258 <220> FEATURE:  
259 <221> NAME/KEY: MOD\_RES  
260 <222> LOCATION: (1)..(1)  
261 <223> OTHER INFORMATION: AcHNCH<sub>2</sub>CH<sub>2</sub>N(CH<sub>2</sub>CH<sub>2</sub>)<sub>2</sub>NCH<sub>2</sub>C(=O)-proline  
264 <400> SEQUENCE: 12  
wt 266 Xaa Leu Gly Leu Leu  
267 1 5  
269 <210> SEQ ID NO: 13  
270 <211> LENGTH: 5  
271 <212> TYPE: PRT  
272 <213> ORGANISM: Artificial  
274 <220> FEATURE:  
275 <223> OTHER INFORMATION: no comment  
277 <220> FEATURE:  
278 <221> NAME/KEY: MOD\_RES  
279 <222> LOCATION: (1)..(1)  
280 <223> OTHER INFORMATION: AcN(CH<sub>2</sub>CH<sub>2</sub>)<sub>2</sub>NCH<sub>2</sub>C(=O)-proline  
283 <400> SEQUENCE: 13  
wt 285 Xaa Leu Gly Leu Leu

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/808,832

DATE: 03/30/2001

TIME: 15:15:26

Input Set : A:\PTO.txt

Output Set: N:\CRF3\03302001\I808832.raw

286 1 5  
288 <210> SEQ ID NO: 14  
289 <211> LENGTH: 4  
290 <212> TYPE: PRT  
291 <213> ORGANISM: Artificial  
293 <220> FEATURE:  
294 <223> OTHER INFORMATION: no comment  
296 <220> FEATURE:  
297 <221> NAME/KEY: MOD\_RES  
298 <222> LOCATION: (4)..(4)  
299 <223> OTHER INFORMATION: O-benzyl-serine  
302 <400> SEQUENCE: 14  
OK 304 Pro Leu Gly Xaa  
305 1  
307 <210> SEQ ID NO: 15  
308 <211> LENGTH: 4  
309 <212> TYPE: PRT  
310 <213> ORGANISM: Artificial  
312 <220> FEATURE:  
313 <223> OTHER INFORMATION: no comment  
315 <220> FEATURE:  
316 <221> NAME/KEY: MOD\_RES  
317 <222> LOCATION: (1)..(1)  
318 <223> OTHER INFORMATION: acetyl-proline  
OK 321 <400> SEQUENCE: 15  
OK 323 Xaa Leu Gly Leu  
324 1  
326 <210> SEQ ID NO: 16  
327 <211> LENGTH: 5  
328 <212> TYPE: PRT  
329 <213> ORGANISM: Artificial  
331 <220> FEATURE:  
332 <223> OTHER INFORMATION: no comment  
334 <220> FEATURE:  
335 <221> NAME/KEY: MOD\_RES  
336 <222> LOCATION: (1)..(1)  
337 <223> OTHER INFORMATION: acetyl-glycine  
OK 340 <400> SEQUENCE: 16  
OK 342 Xaa Pro Leu Gly Leu  
343 1 5  
345 <210> SEQ ID NO: 17  
346 <211> LENGTH: 6  
347 <212> TYPE: PRT  
348 <213> ORGANISM: Artificial  
350 <220> FEATURE:  
351 <223> OTHER INFORMATION: no comment  
353 <220> FEATURE:  
354 <221> NAME/KEY: MOD\_RES  
355 <222> LOCATION: (1)..(1)

Please correct these errors  
in subsequent sequences,  
too.

FWI  
Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

## VERIFICATION SUMMARY

DATE: 03/30/2001

PATENT APPLICATION: US/09/808,832

TIME: 15:15:27

Input Set : A:\PTO.txt

Output Set: N:\CRF3\03302001\I808832.raw

L:10 M:270 C: Current Application Number differs, Replaced Current Application No  
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:39 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1  
L:64 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2  
L:83 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3  
L:102 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4  
L:127 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5  
L:152 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6  
L:177 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7  
L:209 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9  
L:228 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10  
L:247 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11  
L:266 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12  
L:285 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13  
L:304 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14  
L:323 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15  
L:342 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:16  
L:373 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17  
L:398 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18  
L:429 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19  
L:460 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20  
L:491 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21  
L:516 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22  
L:541 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23  
L:566 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24  
L:591 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25  
L:616 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26  
L:641 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27  
L:666 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28  
L:691 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29  
L:710 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30  
L:729 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31  
L:748 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32  
L:773 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33  
L:804 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34  
L:823 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35  
L:842 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36  
L:861 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:37  
L:886 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:38  
L:905 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:39  
L:924 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:40  
L:943 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:41  
L:962 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:42  
L:981 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:43  
L:1000 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:44  
L:1019 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:45  
L:1038 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:46  
L:1057 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:47



VERIFICATION SUMMARY

PATENT APPLICATION: US/09/808,832

DATE: 03/30/2001

TIME: 15:15:27

Input Set : A:\PTO.txt

Output Set: N:\CRF3\03302001\I808832.raw

L:1076 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:48  
L:1095 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:49  
L:1114 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:50  
L:1133 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51